

Figure 1

↙ 30

	32a	32b	32c
	↙	↙	↙
	Atom-Pair Type	Distance Value (Angstroms)	Well-Depth Value (-kcal/mol)
34a ~	CF-CO	4.2	0.1917
34b ~	CF-CN	4.2	0.3797
34c ~	CF-NC	3.55	0.4225
34d ~	CF-NP	4.2	0.3025
34e ~	CP-OA	3	0.4325
34f ~	NC-CO	3.8	1.4381
34g ~	OC-OD	2.6	1.4604
34h ~	OA-ND	2.9	0.7225
34i ~	SA-NR	4.4	0.6577

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Figure 2

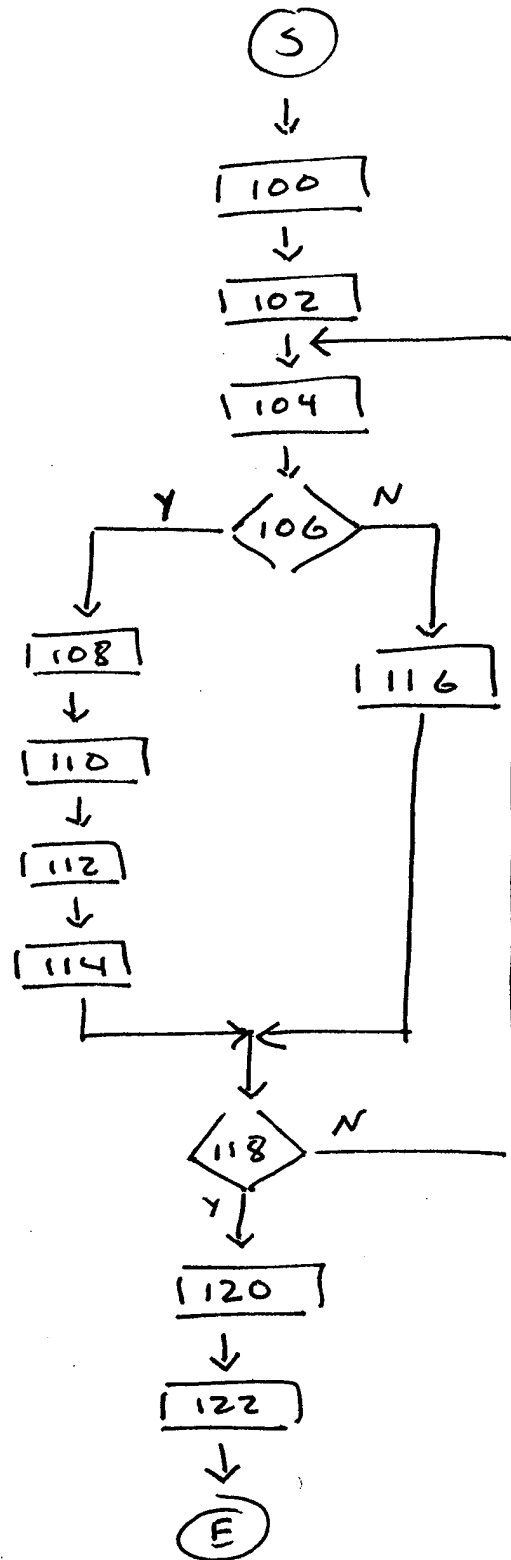


Figure 3

100. USER SPECIFIES PROTEIN-LIGAND COMPLEX
102. PMF-SCORING MODULE ACCESSES PMF-SCORING DATA ASSOCIATED WITH SPECIFIED PROTEIN-LIGAND COMPLEX
104. PMF-SCORING MODULE IDENTIFIES PROTEIN-LIGAND ATOM PAIR IN SPECIFIED PROTEIN-LIGAND COMPLEX
106. SHOULD REPULSION TERM BE USED TO CALCULATE PMF OF IDENTIFIED PROTEIN-LIGAND ATOM PAIR?
108. PMF-SCORING MODULE ACCESSES TABLE OF EMPIRICALLY DERIVED MINIMUM BINDING-ENERGY DISTANCE AND WELL-DEPTH VALUES
110. PMF-SCORING MODULE USES ACCESSED TABLE TO DETERMINE A MINIMUM BINDING-ENERGY DISTANCE VALUE AND A WELL-DEPTH VALUE THAT CORRESPOND TO IDENTIFIED PROTEIN-LIGAND ATOM PAIR
112. PMF-SCORING MODULE USES DETERMINED MINIMUM BINDING-ENERGY DISTANCE AND WELL-DEPTH VALUES TO CALCULATE REPULSION TERM
114. PMF SCORING MODULE USES CALCULATED REPULSION TERM TO CALCULATE PMF OF IDENTIFIED PROTEIN-LIGAND ATOM PAIR
116. PMF SCORING MODULE CALCULATES PMF OF IDENTIFIED PROTEIN-LIGAND ATOM PAIR WITHOUT REPULSION TERM
118. PMF CALCULATED FOR EVERY PROTEIN-LIGAND ATOM PAIR IN SPECIFIED PROTEIN-LIGAND COMPLEX?
120. PMF-SCORING MODULE USES CALCULATED PMFs TO CALCULATE PMF SCORE OF SPECIFIED PROTEIN-LIGAND COMPLEX
122. PMF-SCORING MODULE COMMUNICATES CALCULATED PMF SCORE TO USER

Figure 3 (continued)